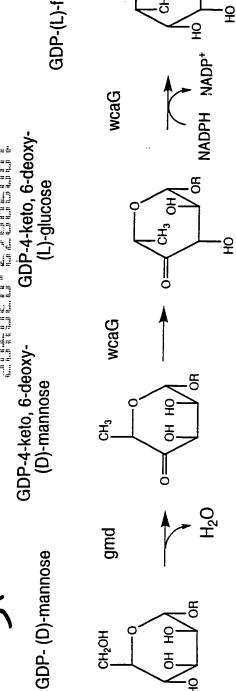
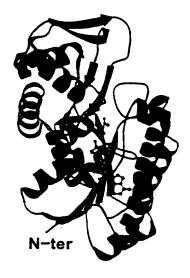
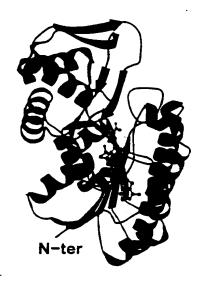
GDP-(L)-fucose

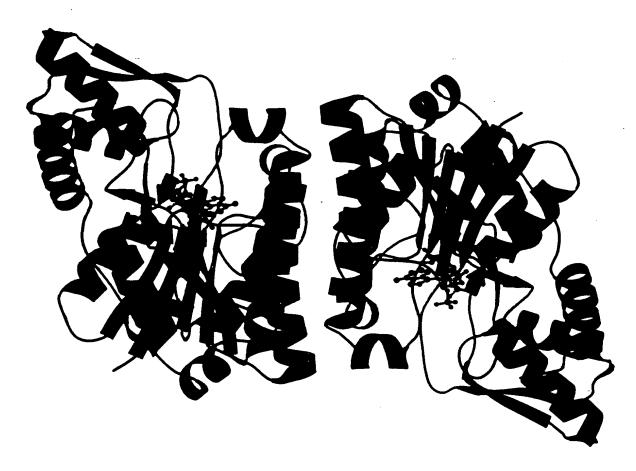
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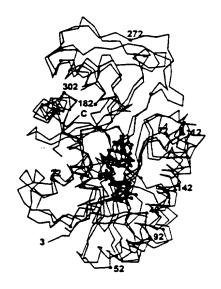


R-GDP









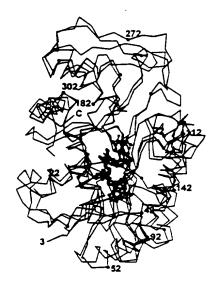
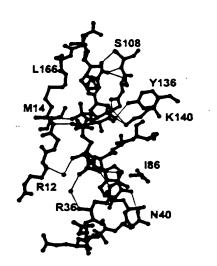
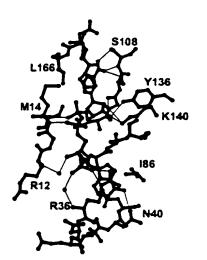
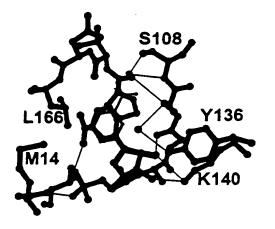
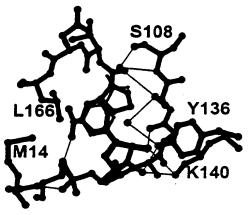


Figure 4 galE coli\_GFS human\_GFS consensus 59 IRNEALMTEILHDHAIDTVIHFAGLKA-VGESVQKPLEYYDNNVNGTLRLISAMRAANVK 41 LLDSRAVHDFFASERIDQVYLAAKVGGIVANNTYPADSIYQNNMIESNIIHAAHQNDVN 48 LTDTAQTRALPEKVQPTHVIHLAAMVGGLPRNIKYNDDFWRKNVHMNDNVLHSAPEVGAR galE coli\_GFS human\_GFS consensus galE coli\_GFS human\_GFS consensus 174 LERYENPYGAEPSGDMGEDPOGIPNMENEYEAQVAVGR----RDSEAFFGNDYPTEDGTG
160 SVMPTMLYGPH-----DNFHPSNSHVIPALERBHEATAQNAPDVVVWGS-----GTP
167 AVIPTMVFGPHDNF-----NIEDGHVLPGLIEKVHLAKS-SGSALTVWGTGY----NE galE coli\_GFS human\_GFS consensus 230 VRDYHUUDLADGEVVAMEKLANK-----PGVHTYNLGAGVGNSVLDVVNAFSKACGKP 208 WREPLHVDDWAAASIHVMELAHEVWLENTQPMLSHINVCTGVDCTIRELAQTIAKVVGYK 214 REQPIYSLDLAQLFHWVLREYNE-----VEPILLSVCEEDEVSIKEAAEAVVEAMDEH galE ČÖli\_GFS Muman\_GFS consensus 241 .\*..... 284 VNYHEAPRREGDLEAYWADASKADREINWRVTRELDEMAQDIWHWQSRHPQGYPD 268 GRVVPDASKPDGTPRKLIDVTRIHQ-IGWYHEISLEAGLASTYQWFLEMQDRE 267 GEVTPDTTKSDGOPKKTASNSKLRTYLPDERFTPFKQAVKETCAWFTDNYEQARK gals coli\_GFS human\_GFS consensus 301 6000 11A1278



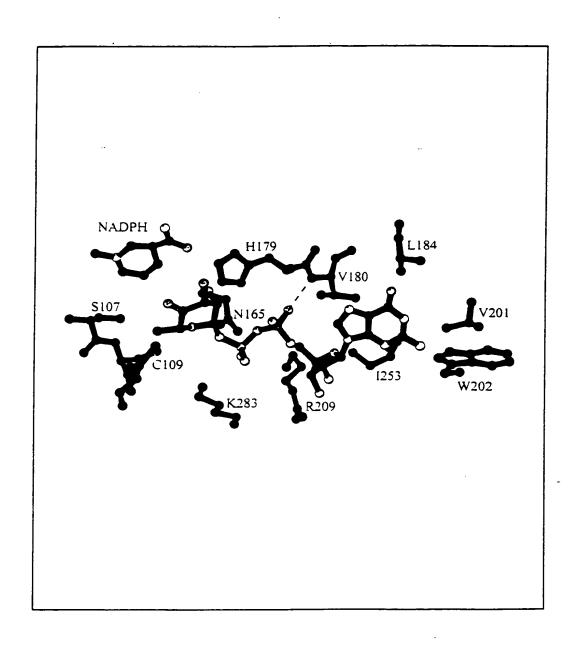






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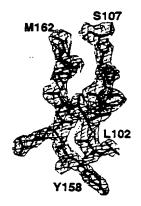
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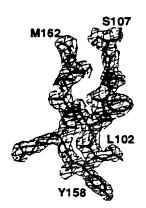
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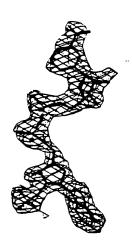




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